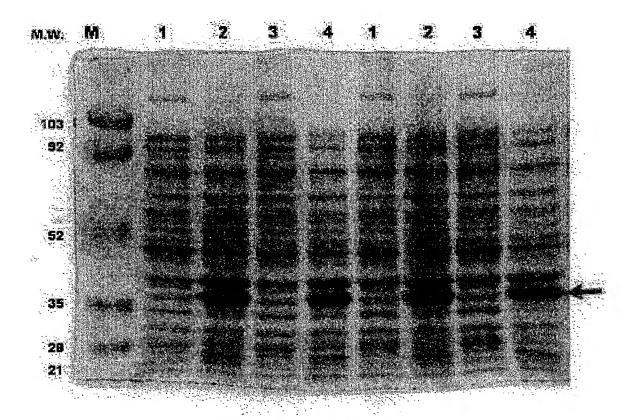
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FIG. 1

At5g10410	MEGTGVVAVYGNGAITEAK-KSPFSVKVGLAQMLRGGVIMDVVNAEQARIAEE 52
At2g38230	MAGTGVVAVYGEGAMTETKQKSPFSVKVGLAQMLRGGVIMDVVNAEQARIAEE 53
At3g16O50	MADQAMTDQDQGAVTLYSGTAITDAKKNHPFSVKVGLAQVLRGGAIVEVSSVNQAKLAES 60
snz3	MSBFKVKTGLAQMLKGGVIMDVVTPEQAIIAER 33
snz2	MSEFKVKTGLAQMLKGGVIMDVVTPEQAIIAER 33
snz1	MTGEDFKIKSGLAQMLKGGVIMDVVTPEQAKIAEK 35
	* * : * * * * : * * : * : * : * : * : *
At5g10410	AGACAVMALERVPADIRAQGGVARMSDPQMIKEIKQAVTIPVMAKARIGHFVEAQILEAI 112
At2g38230	AGACAVMALERVPADIRAQGGVARMSDPEMIKEIKNAVTIPVMAKARIGHFVEAQILEAI 113
At3g16050	AGACSVIVSDPVRSRGGVRRMPDPVLIKEVKRAVSVPVMARARVGHFVEAQILESL 116
snz3	AGACAVMALERIPADMRKSGQVCRMSDPRMIKEIMEAVSIPVMAKVRIGHFVEAQILEEL 93
snz2	AGACAVMALERIPADMRKSGQVCRMSDPRMIKEIMEAVSIPVMAKVRIGHFVEAGILEEL 93
snz1	SGACAVMALESIPADMRKSGKVCRMSDPKMIKDIMNSVSIPVMAKVRIGHFVEAQILEEL 95
SHZI	:***:::: :
At5g10410	GIDYIDESEVLTLADEDHHINKHNFRIPFVCGCRNLGEALRRIREGAAMIRTKG-EAGTG 171
At2g38230	
_	GVDYVDESEVLTLADEDNHINKHNFKIPFVCGCRNLGEALRRIREGAAMIRTKG-EAGTG 172
At3g16050	AVDYIDESEIISVADDDHFINKHNFRSPFICGCRDTGEALRRIREGAAMIRIQGDLTATG 176
snz3	QVDYIDESEVLTPADWTHHIEKHNFKVPFVCGAKDLGEALRRINEGAAMIRTKG-EAGTG 152
snz2	QVDYIDESEVLTPADWTHHIEKHNFKVPFVCGAKDLGEALRRINEGAAMIRTKG-EAGTG 152
snz1	EVDYIDESEVLTPADWTHHIEKDKFKVPFVCGAKDLGEALRRINEGAAMIRTKG-EAGTG 154
	:**:**** ::
At5g10410	NITEAUNUUNGUNGNINU NU NDDDEUEEEAVVI AANUNI UNOEVOI GDI NUUGHAAG OOO
	NIIEAVRHVRSVNGDIRVLRNMDDDEVFTFAKKLAAPYDLVMQTKQLGRLPVVQFAAG 229
At2g38230	NVVEAVRHVRSVNGAIRLLRSMDDDEVFTYAKKIAAPYDLVVQTKELGRLPVVQFAAG 230
At3g16050	NIAETVKNVRSLMGEVRVLNNMDDDEVFTFAKKISAPYDLVAQTKQMGRVPVVQFASG 234
snz3	DVSEAVKHITKIKAEIQQYKENLKTESDFAAKATELRVPVDLLKTTLSEGKLPVVNFAAG 212
snz2	DVSEAVKHITKIKAEIQQYKENLKTESDFAAKATELRVPVDLLKTTLSEGKLPVVNFAAG 212
snz1	DVSEAVKHIRRITEEIKACQQ-LKSEDDIAKVAEEMRVPVSLLKDVLEKGKLPVVNFAAG 213
	:: *:*::: : :: :: :: :: :: :: :: :: :: :
315 40440	
At5g10410	GVATPADAALMMQLGCDGVFVGSGIFKSGDPARRARAIVQAVTHYSDPEMLVEVSCGLGE 289
At2g38230	GVATPADAALMMQLGCDGVFVGSGVFKSGDPVKRAKAIVQAVTNYRDAAVLAEVSCGLGE 290
At3g16050	GITTPADAALMMQLGCDGVFVGSEVFDGPDPFKKLRSIVQAVQHYNDPHVLAEMSSGLEN 294
snz3	GVATPADAALLMQLGCEGVFVGSGIFKSSDPEKLACAIVEATTHYDNPAKLLQVSSDLGD 272
snz2	GVATPADAALLMQLGCEGVFVGSGIFKSSDPEKLACAIVEATTHYDNPAKLLQISSDLGD 272
snz1	GVATPADAALLMQLGCDGVFVGSGIFKSSNPVRLATAVVEATTHFDNPSKLLEVSSDLGE 273
	*::********:***************************
) + E - 1 O 1 1 O	AMUGINI NDENNEDE ANDGE
At5g10410	AMVGINLNDEKVERFANRSE 309
At2g38230	AMVGLNLDD-KVERFASRSE 309
At3g16050	AMESLNVRGDRIQDFGQGSV 314
snz3	LMGGISIQSINEAGGKNGARLSEIGW 298
snz2	LMGGISIQSINEAGGKNGARLSEIGW 298
snz1	LMGGVSIESISHASNGVRLSEIGW 297
	* :::

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FIG. 2



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FIG. 3A

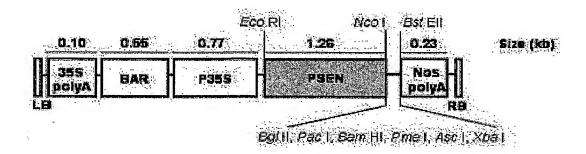


FIG. 3B



FIG. 4



FIG. 5A

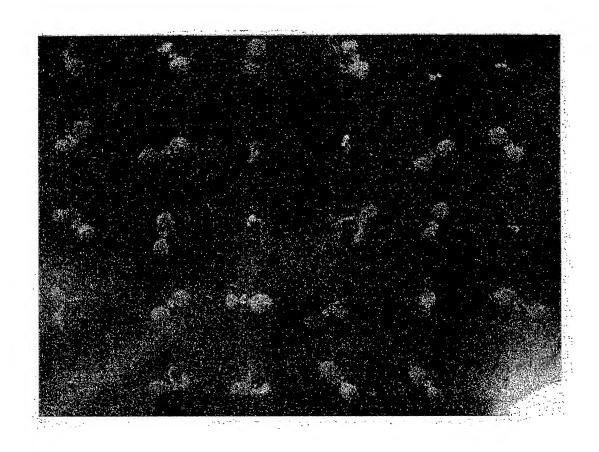
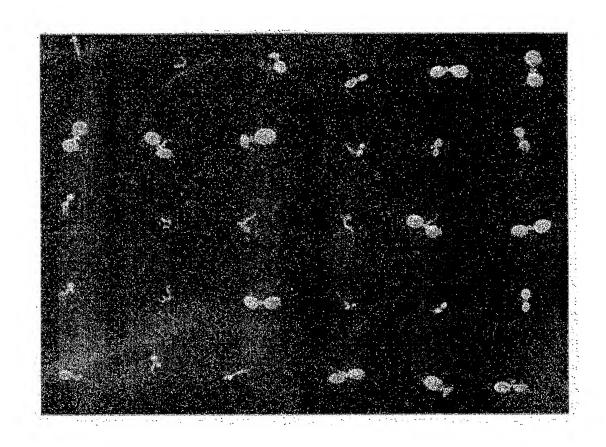
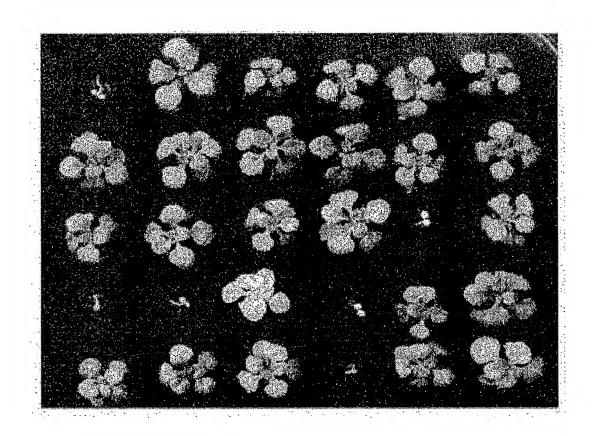


FIG. 5B



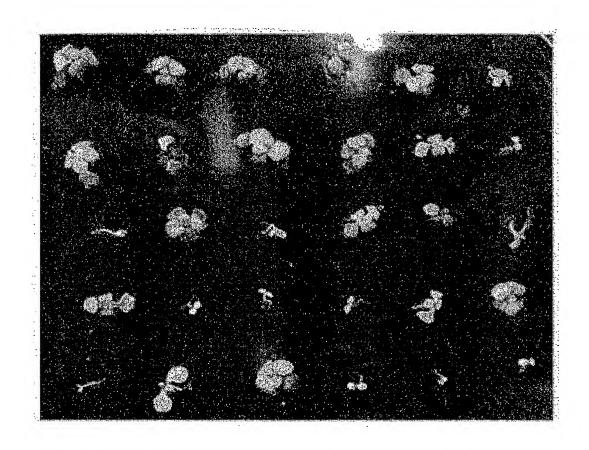
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FIG. 5C



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FIG. 5D



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FIG. 6A

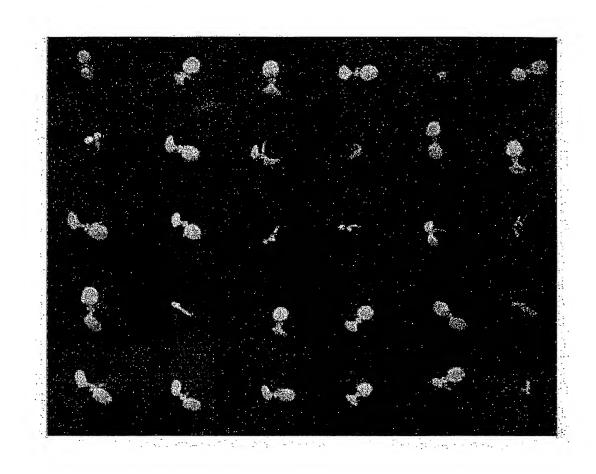
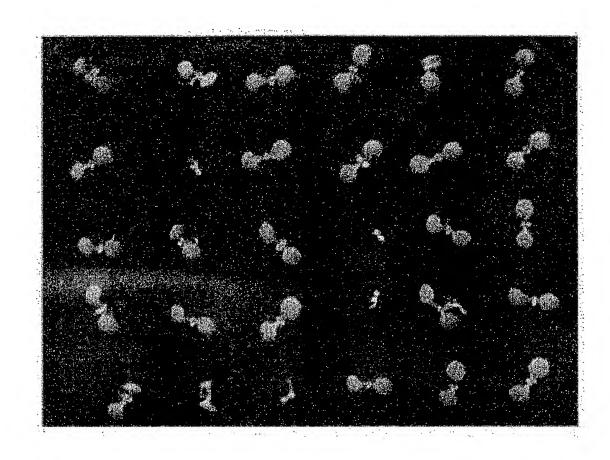
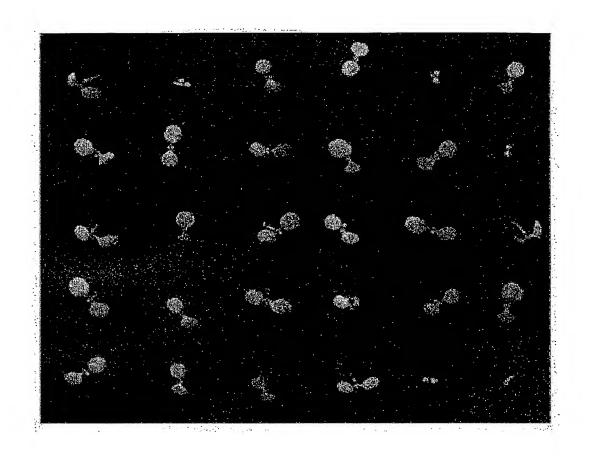


FIG. 6B



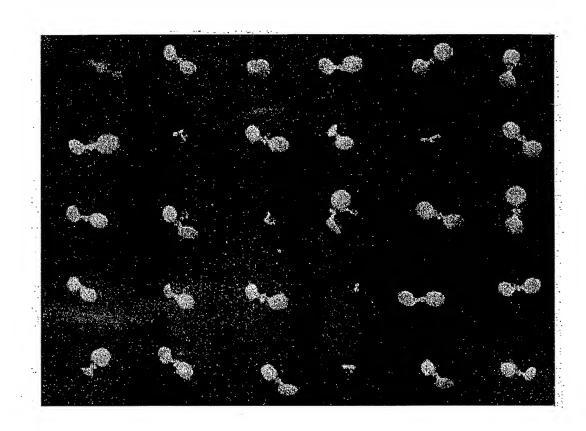
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FIG. 6C



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FIG. 6D



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FIG. 6E

